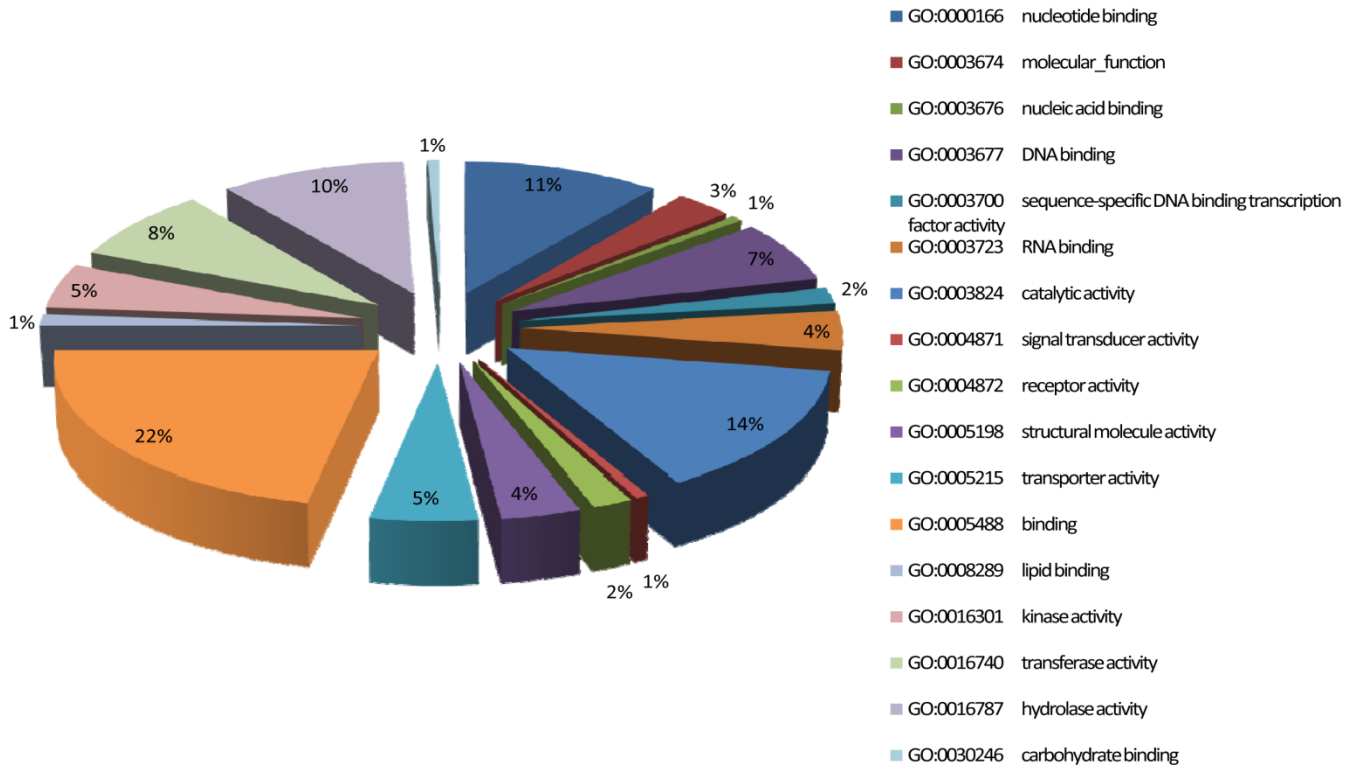
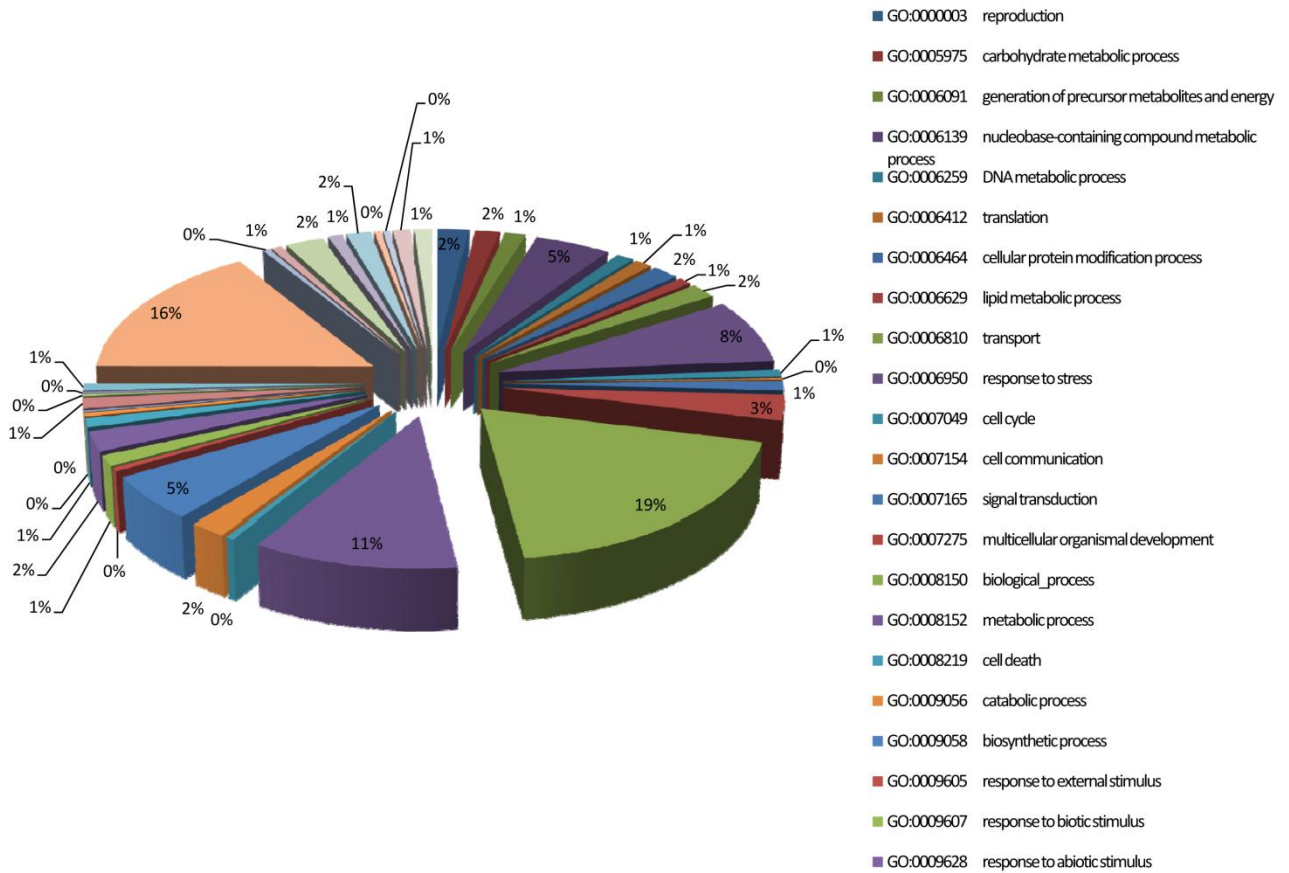


(a)



(b)



(c)

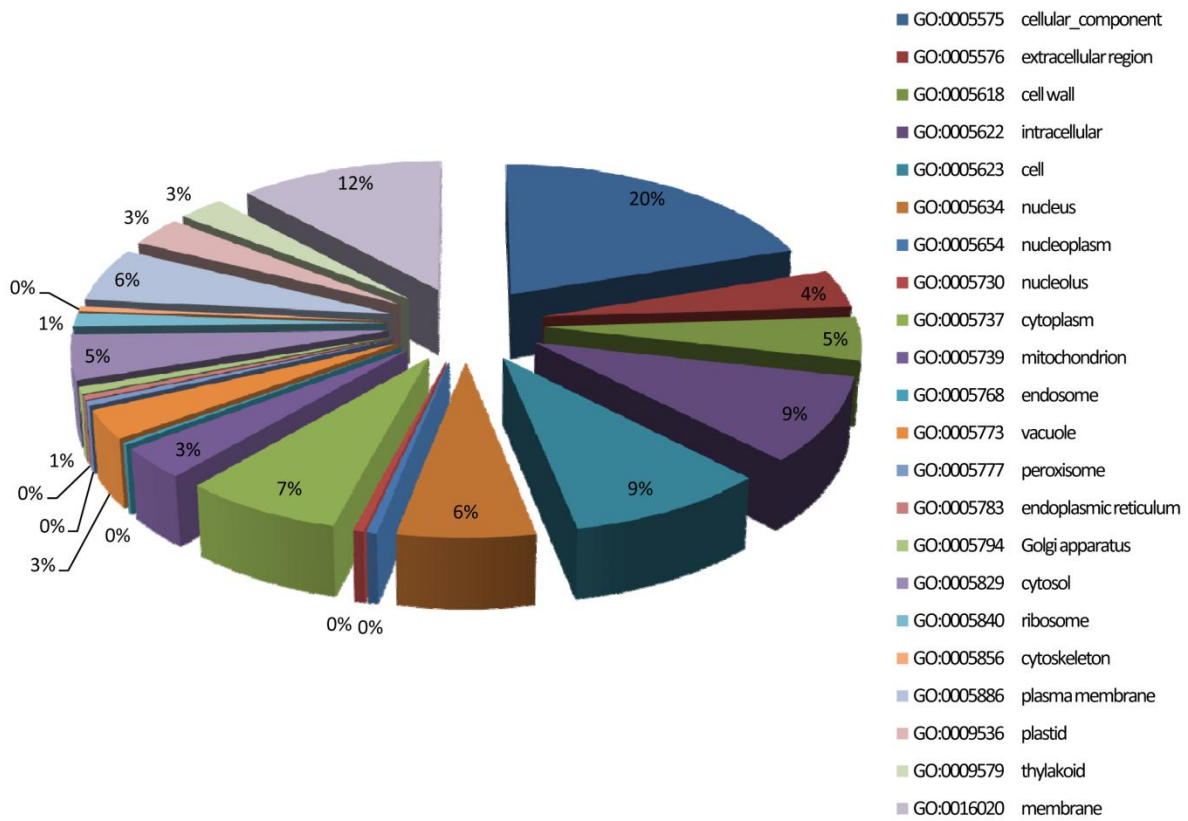


Figure S3. GO-slim annotations of gene products predicted from nine non-neutral candidate genomic regions between fiber flax and linseed groups. **(a)** Molecular function. **(b)** Biological process. **(c)** Cellular component.